

1 ecorI TCTCCATACT TTGGATAAGG AAAATACAGAC ATGAAAAAATC TCATTGCTGA GTTGTATT AAGCTTGCCC AAAAGAAGA AGAGTCGAAT
 CTTAAGTTGA AGAGGTATGA AACCTATTCC TTATGCTG TACTTTAG AGTAACGACT CAACAPATAAA TTGAAACGGG TTTTCTCTCT TCTCAGCTTA
 101 GAACTGTTG CGGAGGTAGA AGCTTGGAG ATTATCGTCA CTGCAATGCT TCGCAATATG GCGCAAAATG ACCAACAGCG GTTGATTGAT CAGGTAGGG
 CTTGACACAC GCGTCCATCT TCGAAACCTC TAATAGCAGT GACGTTACGA AGCGTTATAC CGCGTTTAC CGCGTTTAC TGGTTGTCGC CAACTAACTA GTCCCATCTCC
 201 GGGGGTGTAA CGAGGTAAAG CCCGATGCCA GCATTCCCTGA CGACGATAACG GAGCTGCTGC GCGATTACGT AAAGAAGTTA TTGAAGGCATC CTCGTCAGTA
 CCCGGACAT GCTCCATTTC GGGCTACGGT CGTAAGGACT GCTGCTATGC CTCGACGAGC CGCTAATGCA TTCTCTCAAT AACTTCGTAG GAGCAGTCA
 301 AAAAGTTAAT CTTTCAACA GCTGTCAAA AGTTGCTCAAGG GCCGAGACTT ATAGTCGCTT TGTTTTATT TTTTAATGTA TTGTTAACTA GTACGCAAGT
 TTTCAATTAA GAAAAGTTGT CGACAGTATT TCAAACAGTGC CGGCTCTGAA TATCAGCGAA ACAAAAATAA AAAATTACAT AAACATTGAT CATGGTTCA

 Trp SD xbaI STII SD ATG AAA AAG AAT ATC GCA TTT CTT CTT GCA TCT ATG TTC ATC GAA CGT AGA TAC AAG CAA AAA AGA
 401 TCACGTAAA AGGGTATCTA GAGGTTGAGG TGATTAA TAC TTT TTC TTA TAG CGT AAA GAA GAA CGT AGA TAC AAG CAA AAA AGA
 AGTGCATTTT TCCCATAGAT CTCCAACCTCC ACTAAAA Met Lys Lys Asn Ile Ala Phe Leu Ala Ser Met Phe Val Phe Ser
 1

486 ATT GCT ACA AAT GCC TAT GCA (SEQ ID NO: 13)
 TAA CGA TGT TTA CGG ATA CGT
 17 Ile Ala Thr Asn Ala Tyr Ala (SEQ ID NO: 14)

FIG. 1

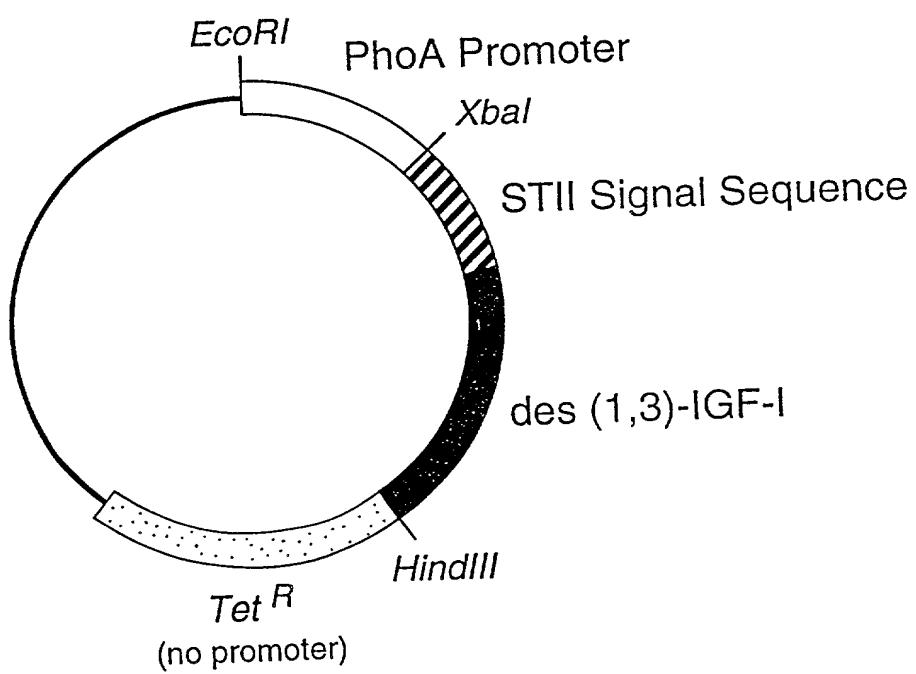
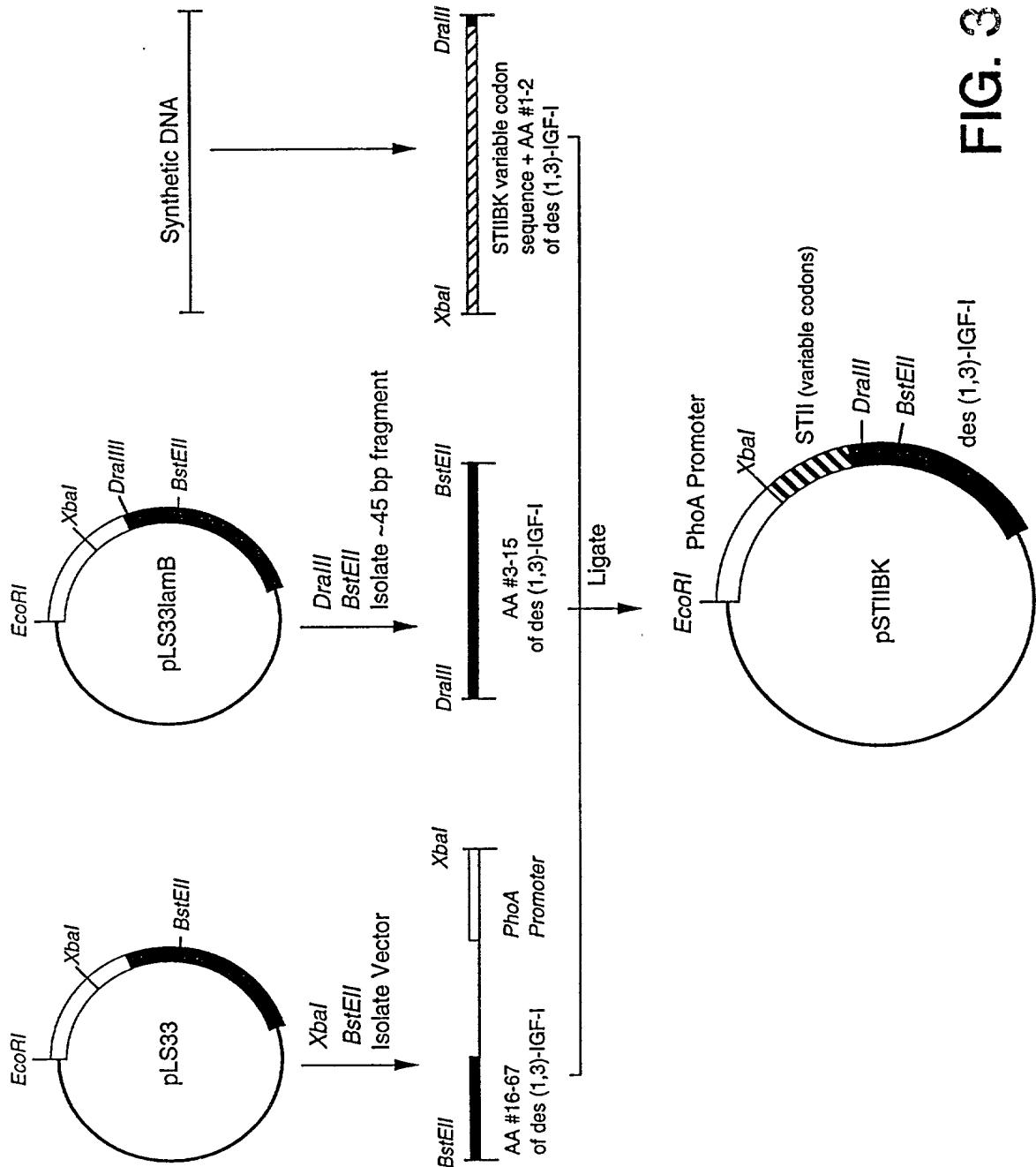


FIG. 2



66

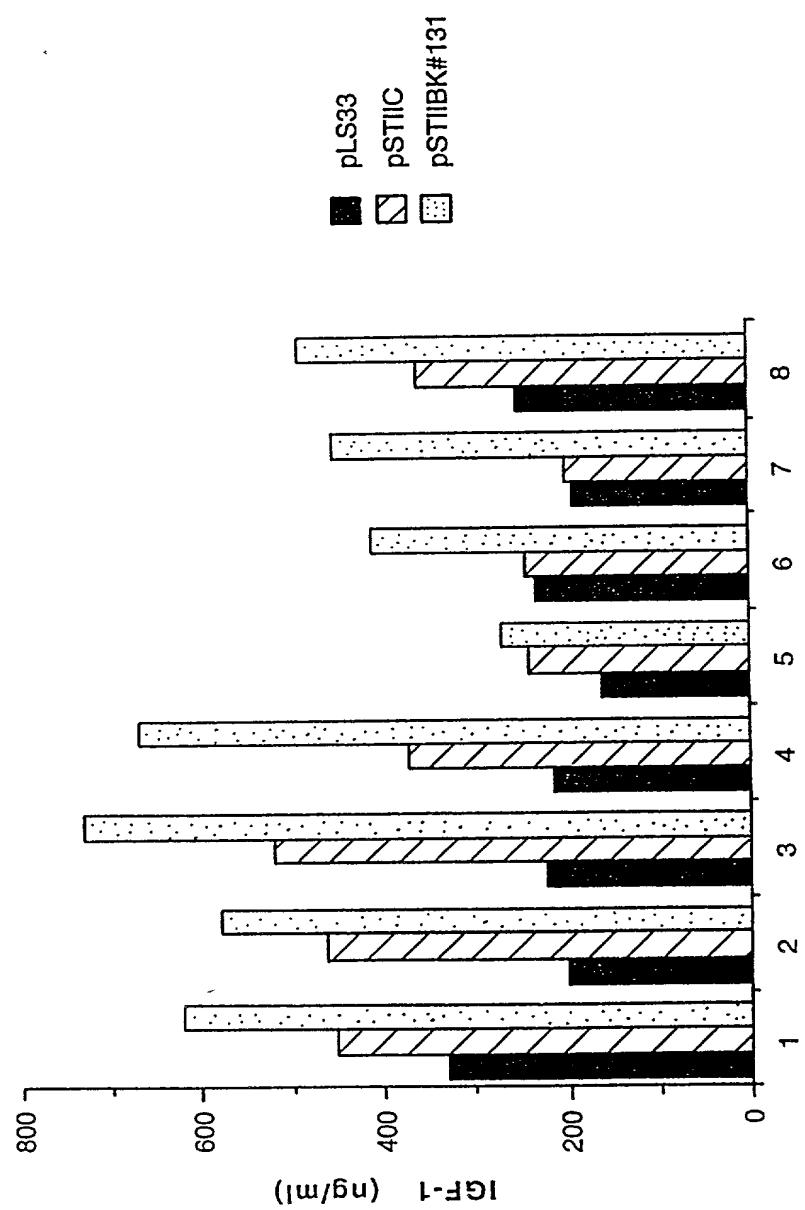


FIG. 4

Experiment

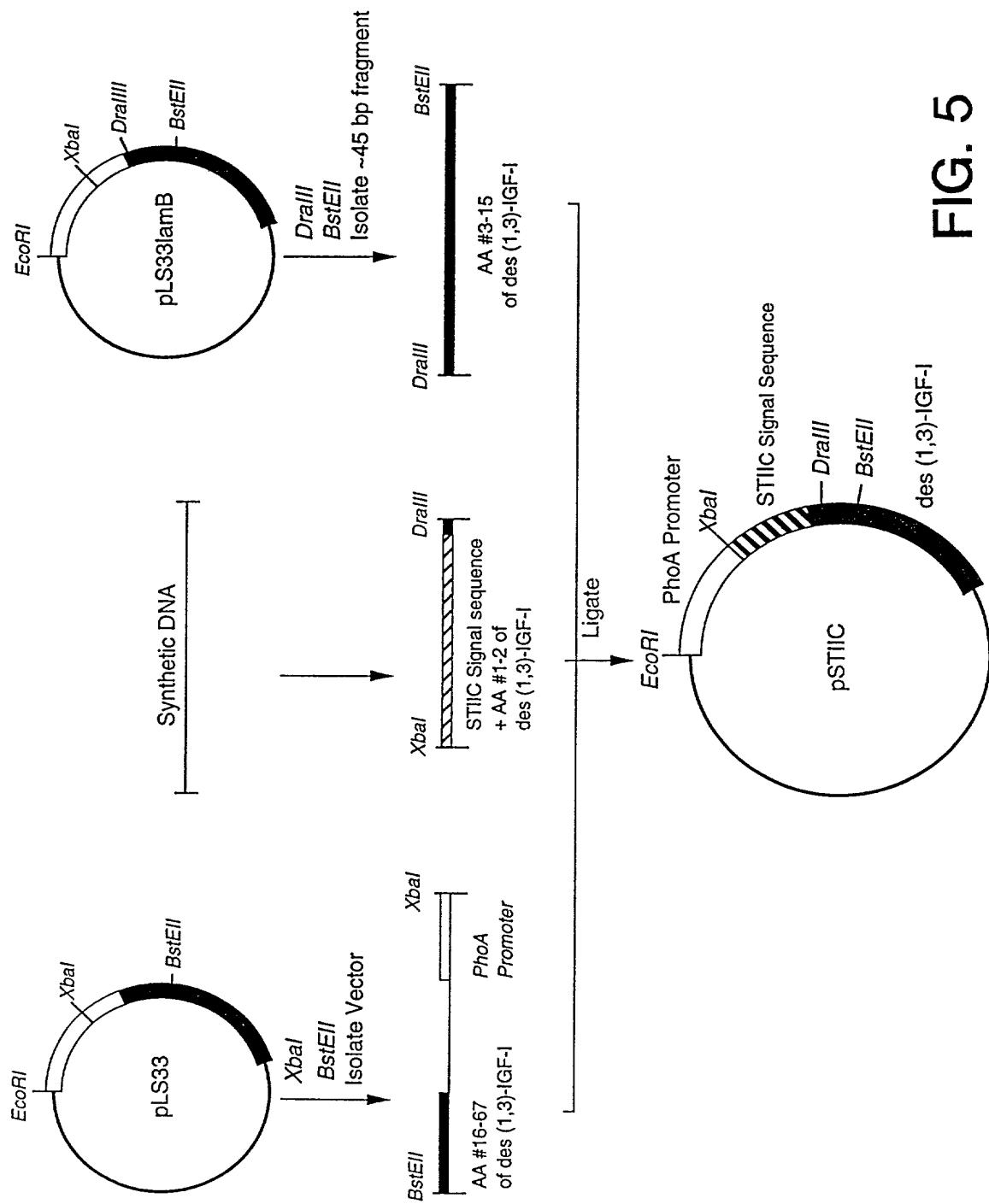


FIG. 5

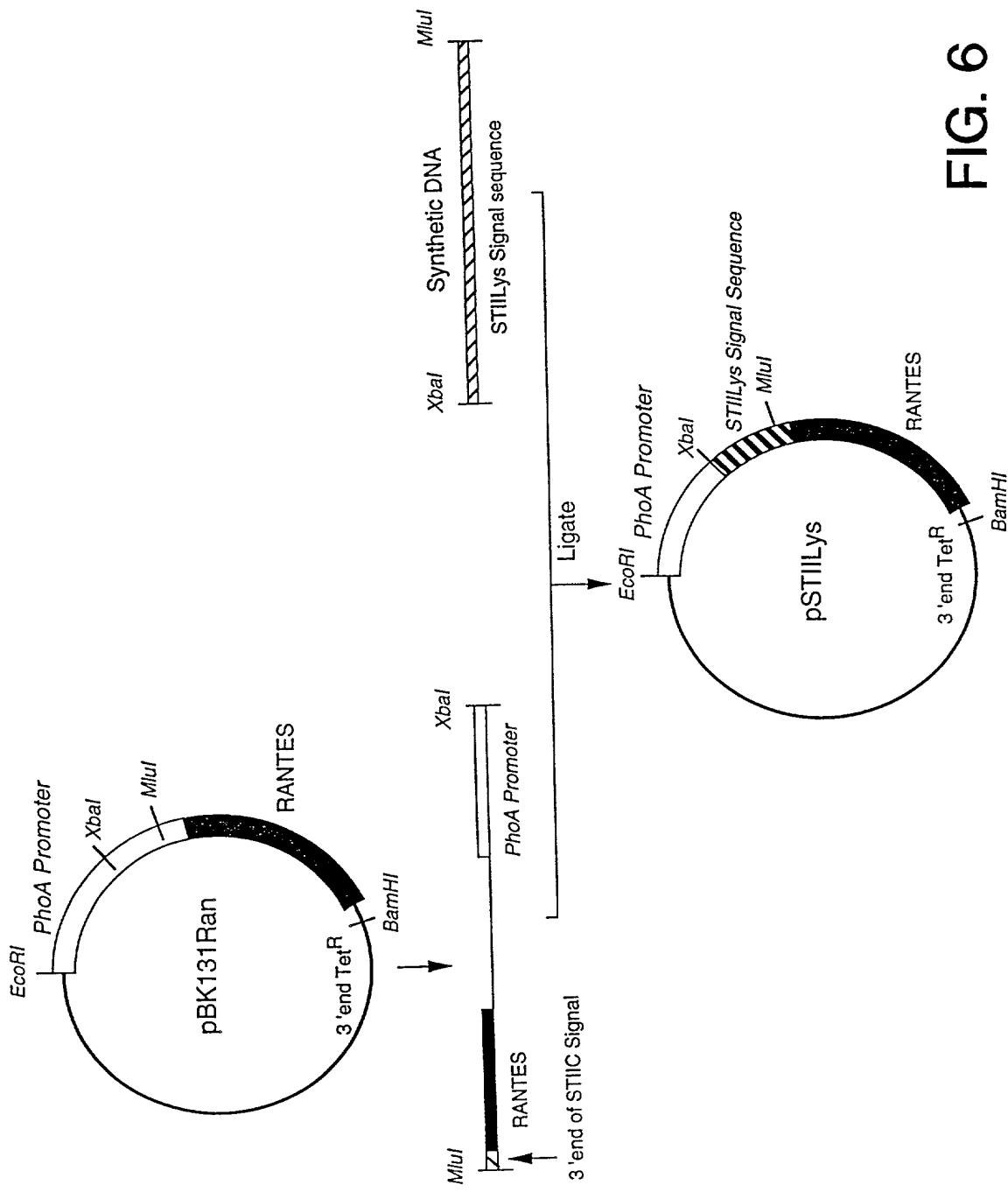


FIG. 6

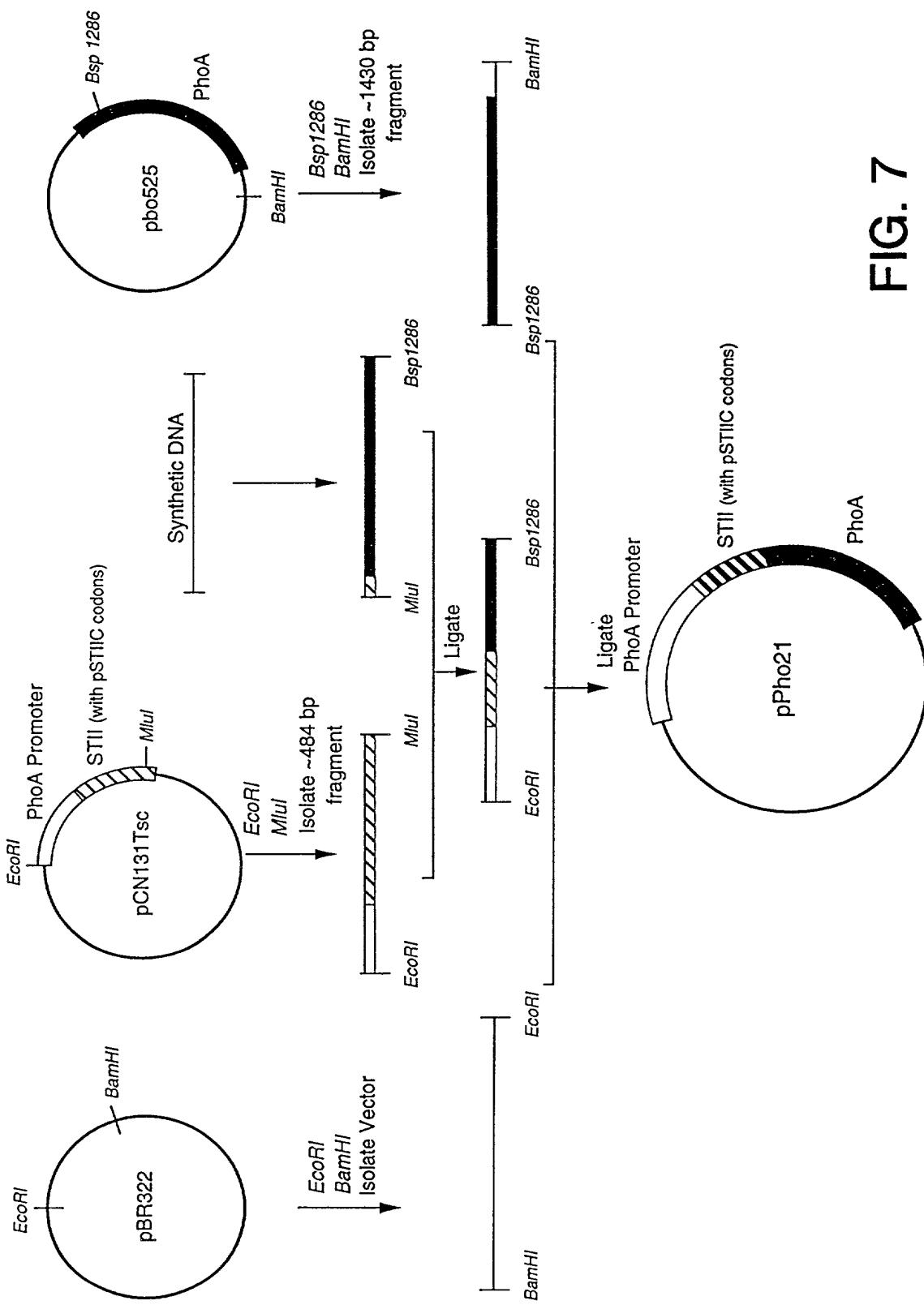


FIG. 7

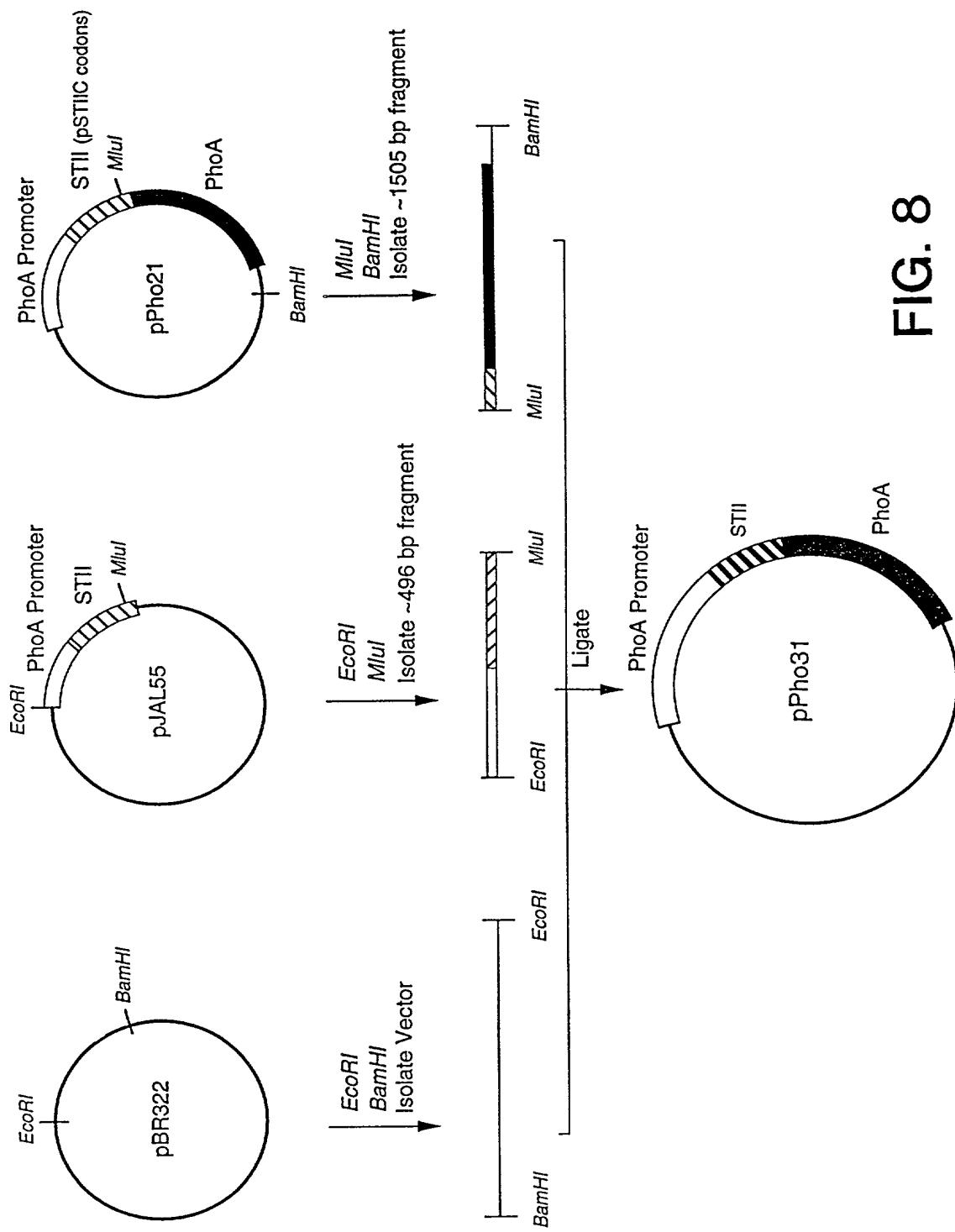


FIG. 8

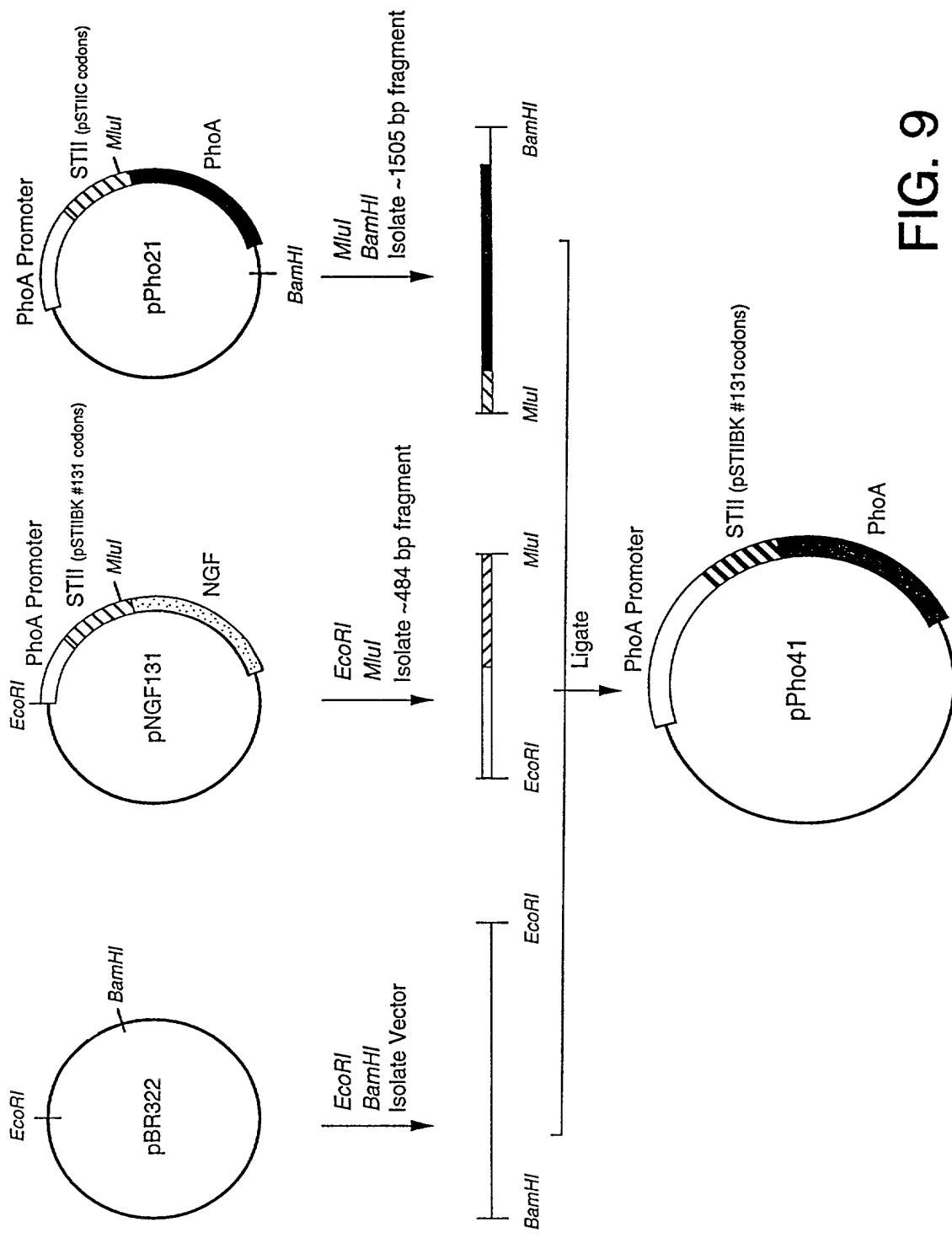


FIG. 9

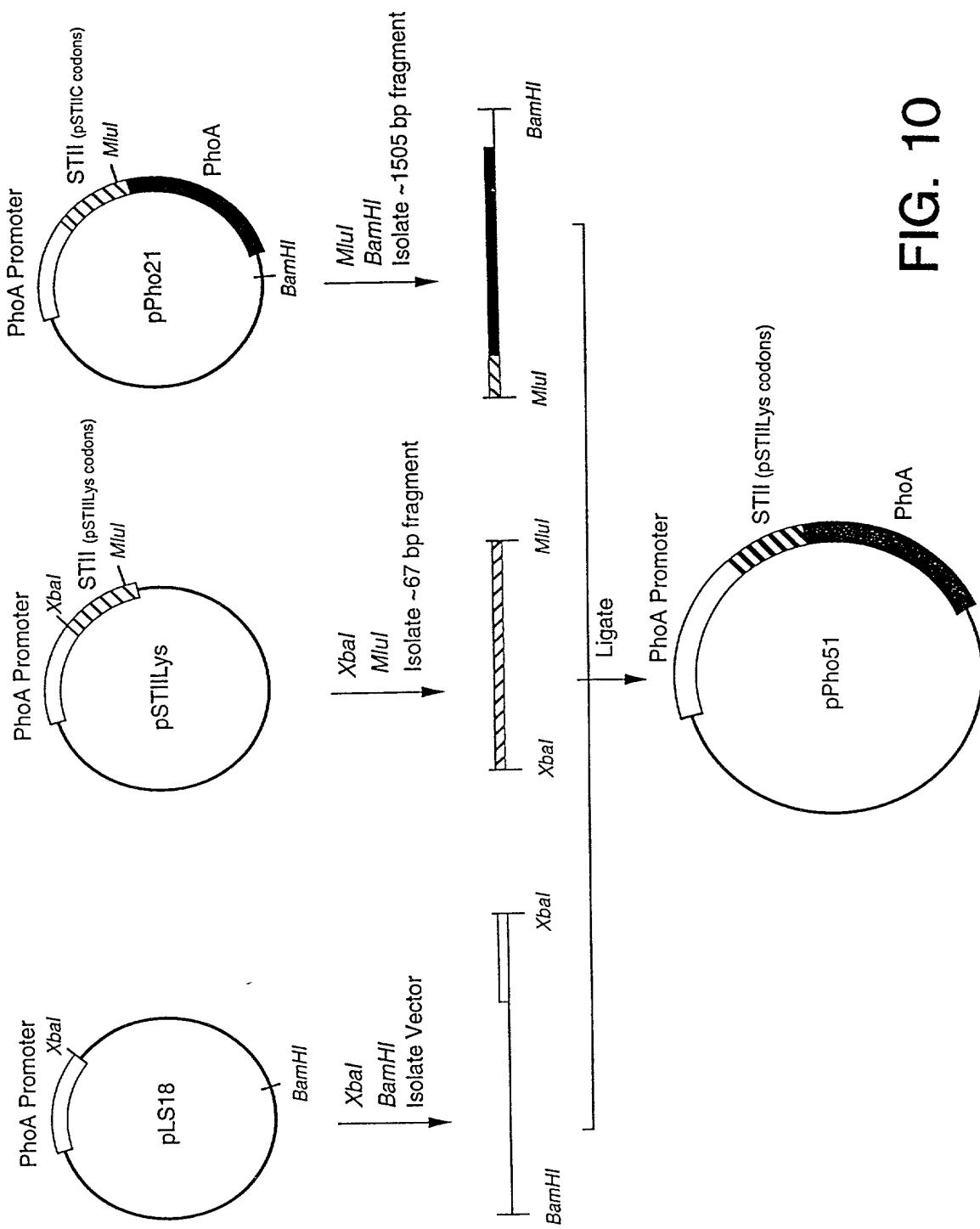


FIG. 10

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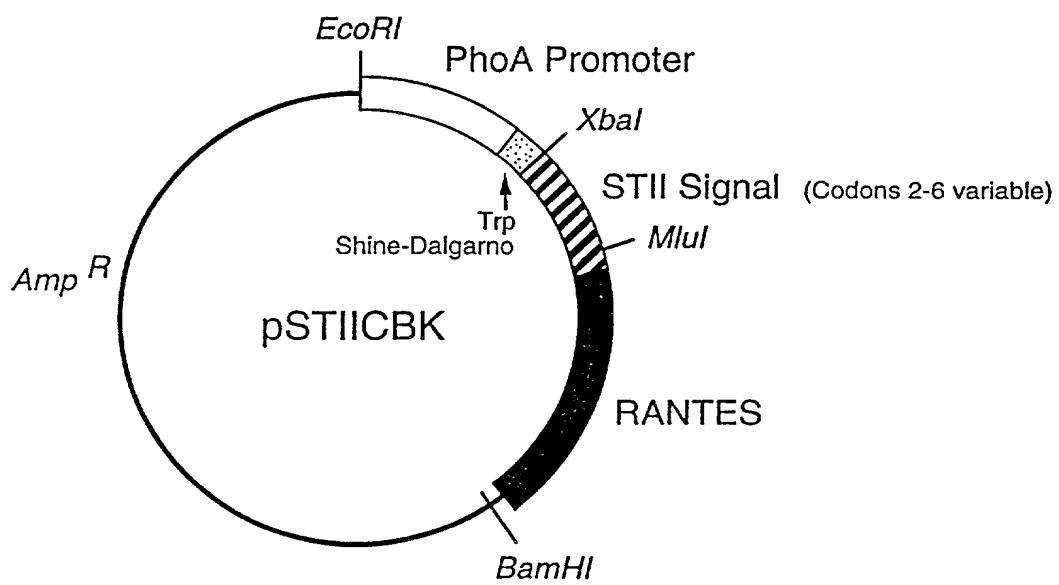


FIG. 11

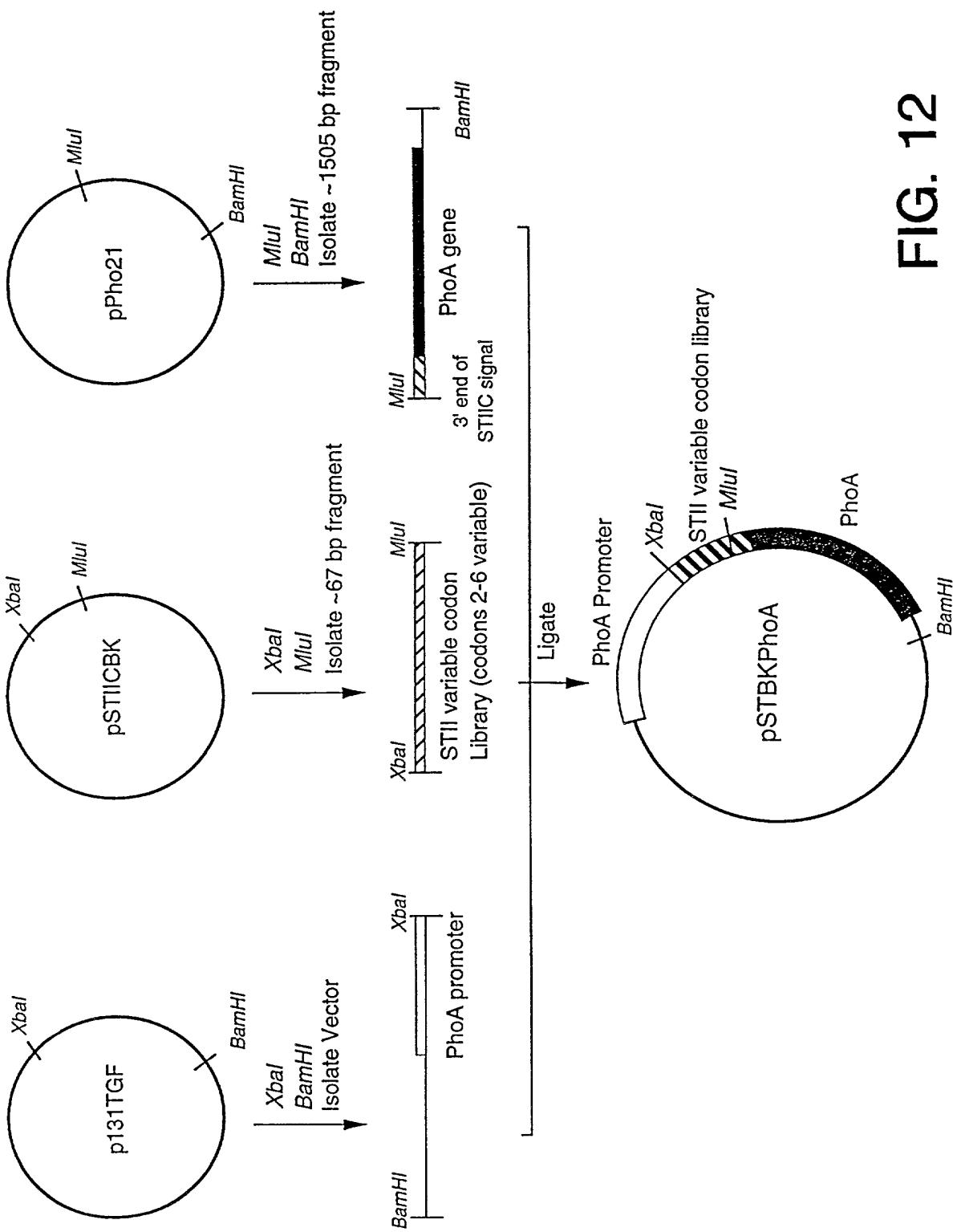


FIG. 12

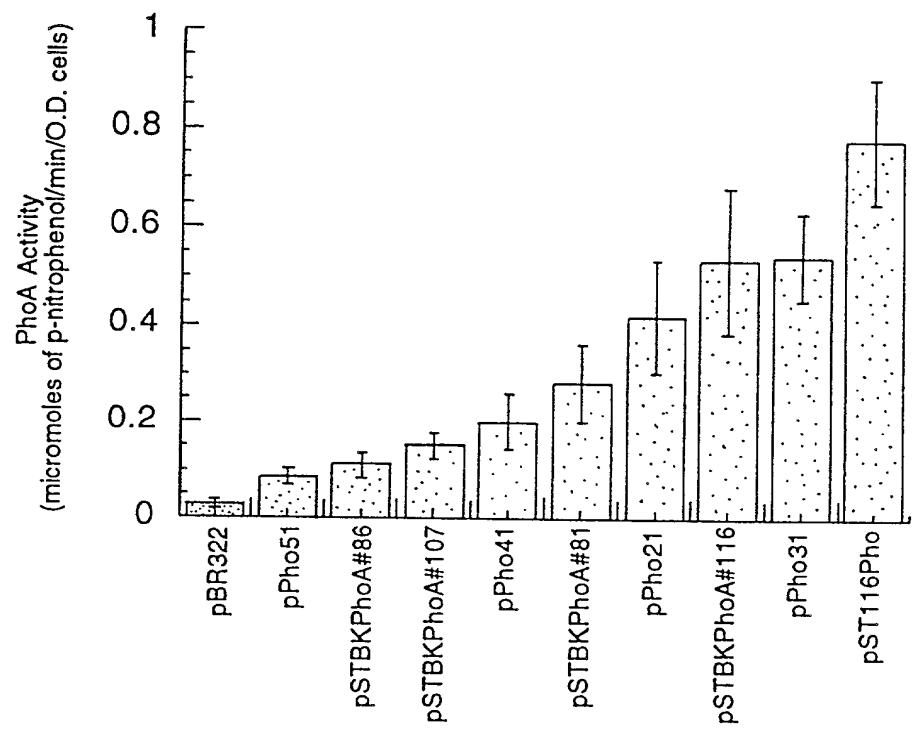


FIG. 13

pPho31 (Wild type STII + Mu1 site)
TCTAGAGTTGAGGTGATTTT ATG AAA AAT ATC GCA TTT CTT GCA TCT ATG TTC GTT

pPho21 (STI IC)
TCTAGAATT ATG AAA AAG AAT ATC GCA TTT CTT GCA TCT ATG TTC GTT

pPho41 (STI IBK#131)
TCTAGAATT ATG AAG AAT ATT GCG TTC CTA CTT GCC TCT ATG TTT GTC

pPho51 (STII Lys - unless otherwise noted this sequence is the TIR=1 used in the examples)
TCTAGAATT ATG AAG AAT ATC GCA TTT CTT GCA TCT ATG TTC GTT

pSTBKPPhoA#116
TCTAGAATT ATG AAA AAC ATC GCA TTT CTT GCA TCT ATG TTC GTT

pSTBKPPhoA#81
TCTAGAATT ATG AAA AAC ATT GCC TTT CTT GCA TCT ATG TTC GTT

pSTBKPPhoA#107
TCTAGAATT ATG AAG AAA AAC ATC GCT TTT CTT GCA TCT ATG TTC GTT

pSTBKPPhoA#86
TCTAGAATT ATG AAA AAG AAC ATA GCG TTT CTT GCA TCT ATG TTC GTT

pST116Pho
TCTAGAGTTGAGGTGATTTT ATG AAA AAC ATC GCA TTT CTT GCA TCT ATG TTC GTT

FIG. 14A

		<u>TIR RELATIVE STRENGTH</u>	
TTT	TCT ATT GCT ACA AAY GCS TAT GCM*	(SEQ ID NO:15)	9
TTT	TCT ATT GCT ACA AAC GCG TAT GCM	(SEQ ID NO:16)	7
TTT	TCT ATA GCT ACA AAC GCG TAT GCM	(SEQ ID NO:17)	3
TTT	TCT ATT GCT ACA AAC GCG TAT GCM	(SEQ ID NO:18)	1
TTT	TCT ATT GCT ACA AAC GCG TAT GCM	(SEQ ID NO:19)	9
TTT	TCT ATT GCT ACA AAC GCG TAT GCM	(SEQ ID NO:20)	4
TTT	TCT ATT GCT ACA AAC GCG TAT GCM	(SEQ ID NO:21)	2
TTT	TCT ATT GCT ACA AAC GCG TAT GCM	(SEQ ID NO:22)	1
TTT	TCT ATT GCT ACA AAC GCG TAT GCM	(SEQ ID NO:23)	13

* The codons for the last four amino acids of this sequence may differ in some of the examples of protein secretion. For example, in the IGF-1, VEGF165 and RANTES secretion plasmids, the sequence is AAT GCC TAT GCA. The last codon for the last amino acid in every sequence listed may vary in the examples of protein secretion - GCC and GCA were both used.

FIG. 14B

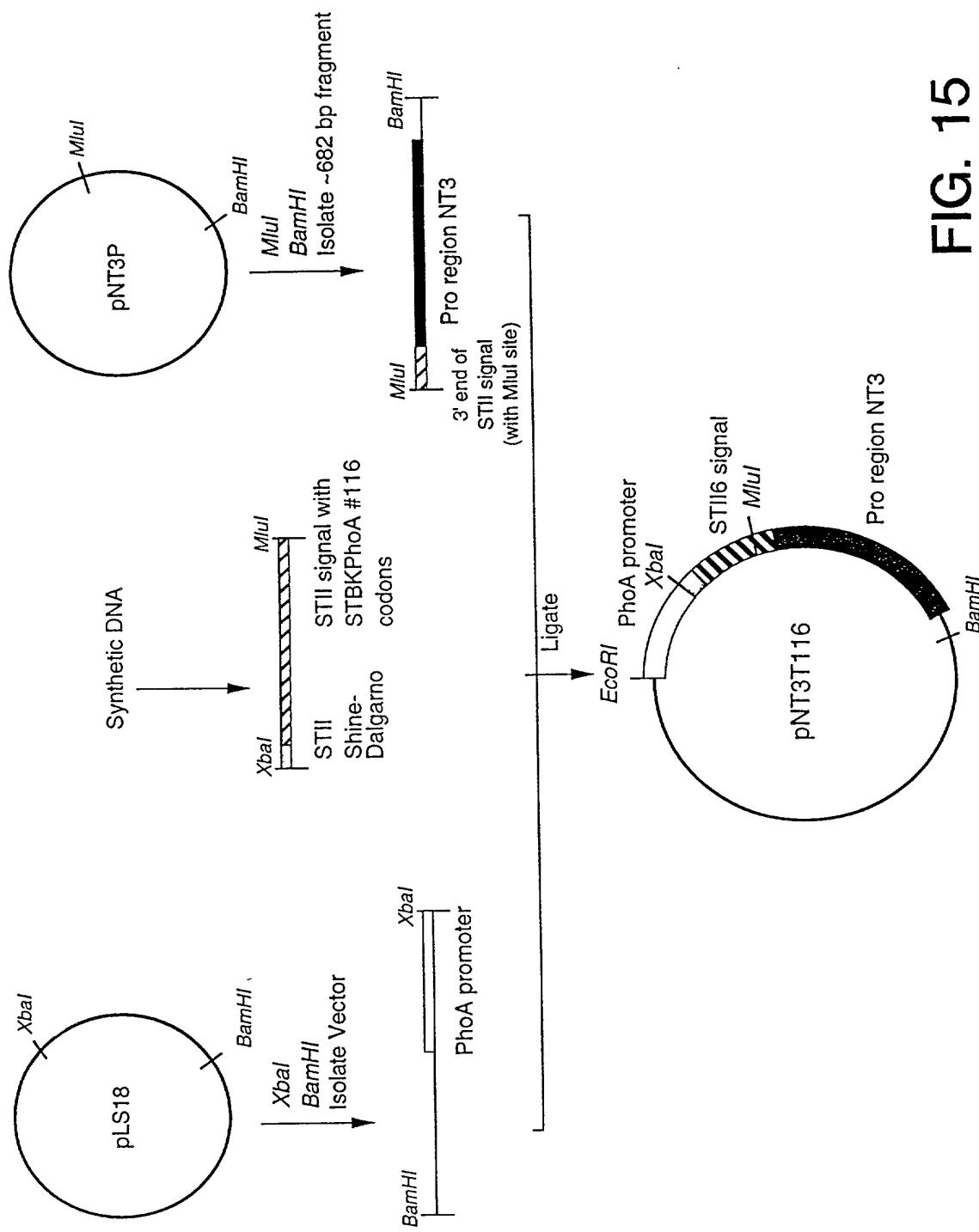


FIG. 15

A-63487-3

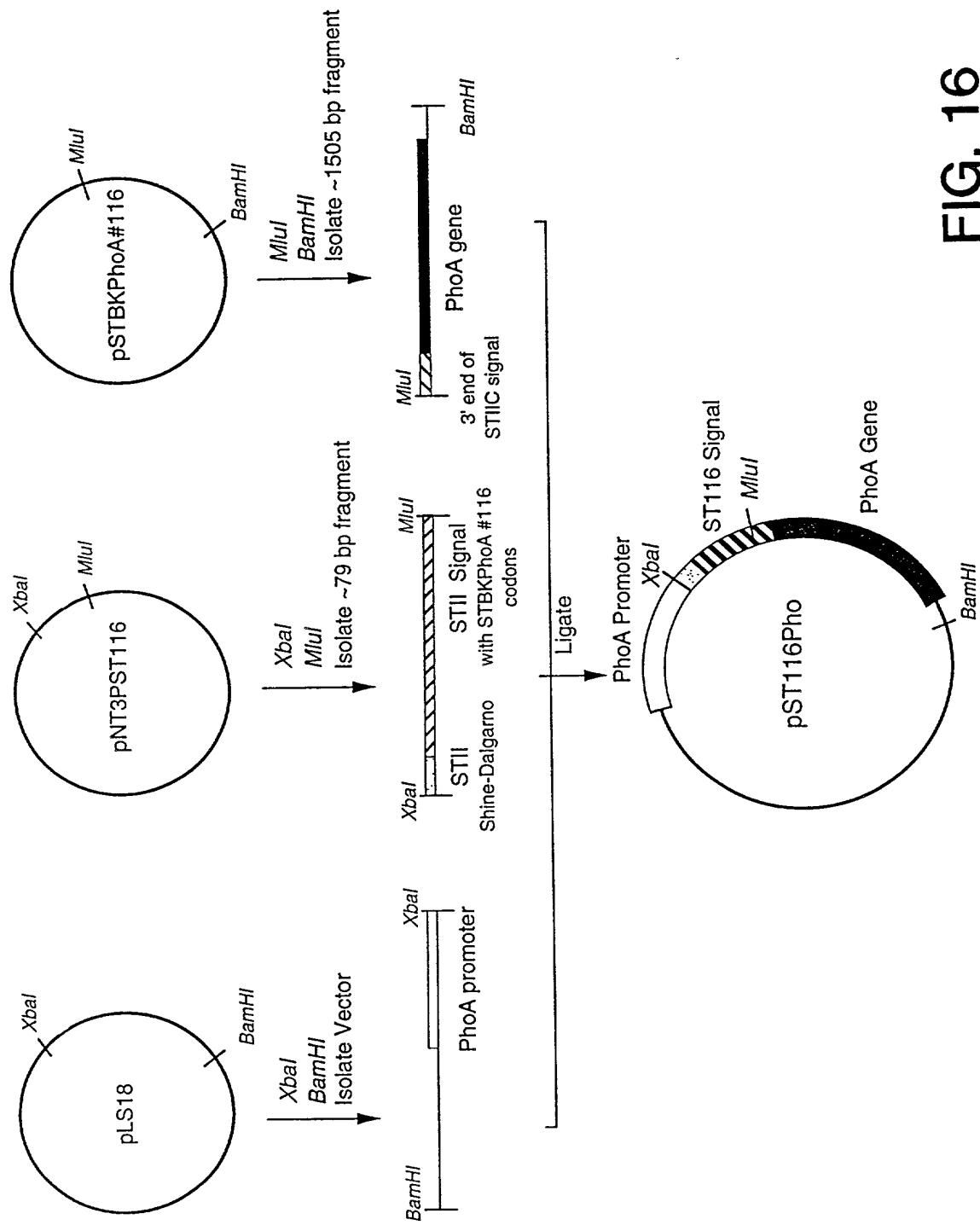
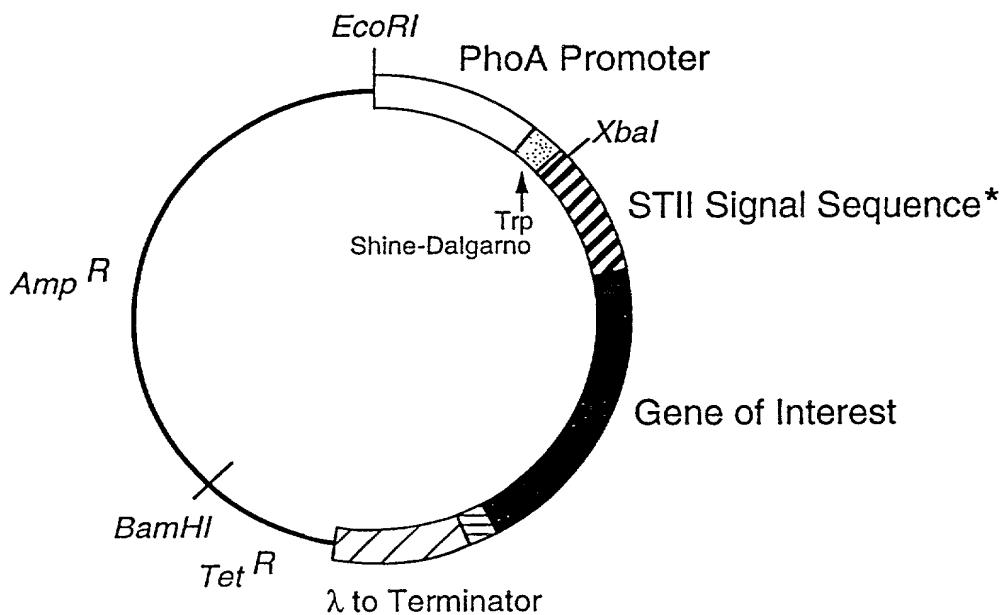
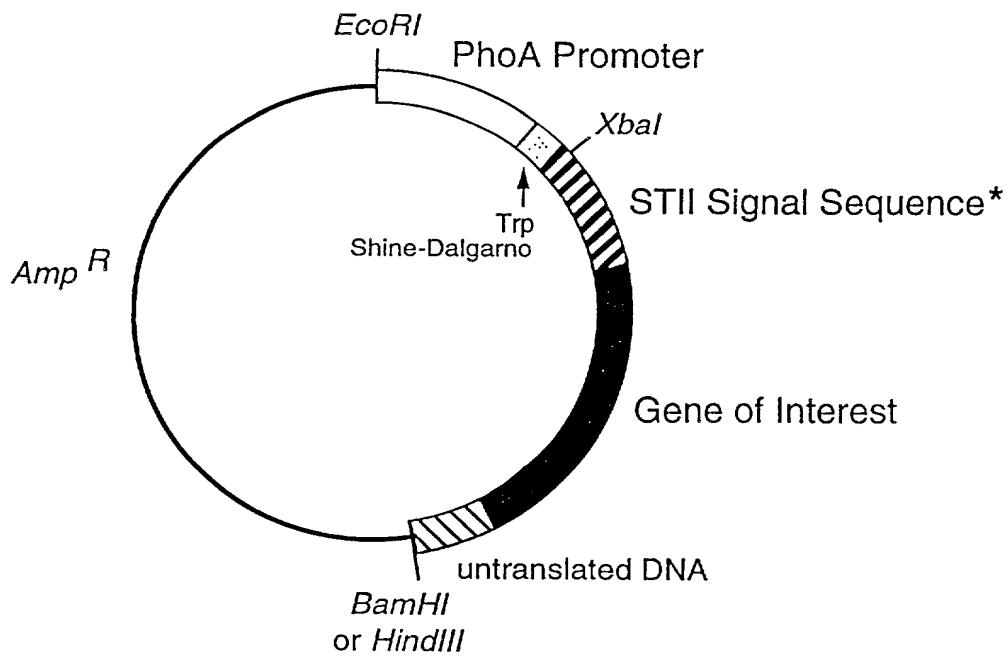


FIG. 16



* One of the nucleotide sequences listed in Figure 14
(STII Shine-Dalgarno may also be included).

FIG. 17



* One of the nucleotide sequences listed in Figure 14
(STII Shine-Dalgarno may also be included).

FIG. 18